```
************************************************************
```

Reviewer Comments:
<210>   6
<211>   6069
<212>   DNA
<213>   Artificial sequence

<220>
<223>   pETM-GFP-Imm plasmid containing Hisx6 tag, flexible joint
as frame adapter, and A. victoria GFP gene

<220>
<221>   CDS
<222>   (1)..(876)
<223>   pETM-GFP-Imm plasmid containing Hisx6 tag, flexible joint
as frame adapter, and A. victoria GFP gene

Two problems above: 1) the first <223> response exceeds the Sequence
Rules' required 72-character line limit; please do not use TAB codes,
and please insert a hard return. 2) please delete the second <223>
line, since it is a duplicate of the first <223> line.

<210>   8
<211>   17
<212>   DNA
<213>   Artificial sequence

<220>
<223>   Frame adapter used for prevention of frameshift mutation as a
    result of plasmid modification

```
<220>
<221>  misc_feature
<222>  (1)..(17)
<223>  Frame adapter used for prevention of frameshift mutation as a
       result of plasmid modification

<400>  8
catgggaggc acggtac                                                    17
```

Please insert a hard return in the first <223> response; don't use TAB
codes. Please delete the second <220>-<223> section since the <223>
response is a duplicate of the first <223> response.

```
<210>  10
<211>  6
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Peptide design based on charge and shape to bind the
expressed protein to a suitably interactive surface



<220>
<221>  MISC_FEATURE
<223>  Peptide design based on charge and shape to bind the
expressed protein to a suitably interactive surface

<400>  10

His His His His His His
1                   5
```

Please insert a hard return in the first <223> response; please remove
the second <220>-<223> section, since it has a duplicate <223> response.

* *

Validated By CRFValidator v 1.0.3

Application No: 10550226   Version No: 3.0

                    Input Set:

                    Output Set:

                    Started: 2009-08-25 12:01:09.452
                   Finished: 2009-08-25 12:01:10.981
                    Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 529 ms
            Total Warnings: 12
              Total Errors: 0
     No. of SeqIDs Defined: 13
         Actual SeqID Count: 13

Error code   Error Description

W   213   Artificial or Unknown found in <213> in SEQ ID (2)

W   213   Artificial or Unknown found in <213> in SEQ ID (3)

W   213   Artificial or Unknown found in <213> in SEQ ID (4)

W   213   Artificial or Unknown found in <213> in SEQ ID (5)

W   213   Artificial or Unknown found in <213> in SEQ ID (6)

W   213   Artificial or Unknown found in <213> in SEQ ID (7)

W   213   Artificial or Unknown found in <213> in SEQ ID (8)

W   213   Artificial or Unknown found in <213> in SEQ ID (9)

W   213   Artificial or Unknown found in <213> in SEQ ID (10)

W   402   Undefined organism found in <213> in SEQ ID (11)

W   213   Artificial or Unknown found in <213> in SEQ ID (12)

W   213   Artificial or Unknown found in <213> in SEQ ID (13)

<110>   SAHIN, ERINC
         TARALP, ALPAY
         SAYERS, SEHRA

<120>   CIRCULAR RECOMBINANT PLASMID DNA CONSTRUCTS AND THEIR PROTEIN
         PRODUCTS, METHODS OF PREPARATION AND IMMOBILISATION OF PROTEINS
         ON SUPPORT

<130>   U015936-2

<140>   10550226
<141>   2006-11-22

<150>   PCT/TR2003/000019
<151>   2003-03-20

<160>   13

<170>   PatentIn version 3.3

<210>   1
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<213>   Aequorea victoria

<220>
<221>   gene
<222>   (17)..(733)
<223>   GFP gene

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tgatgcaaca tacggaaaac ttacccttaa atttatttgc actactggaa aactacctgt   180

tccatggcca acacttgtca ctactttctc ttatggtgtt caatgctttt cccgttatcc   240

ggatcatatg aaacggcatg acttttttcaa gagtgccatg cccgaaggtt atgtacagga   300

acgcactata tctttcaaag atgacgggaa ctacaagacg cgtgctgaag tcaagtttga   360

aggtgatacc cttgttaatc gtatcgagtt aaaaggtatt gattttaaag aagatggaaa   420

cattctcgga cacaaactcg agtacaacta taactcacac aatgtataca tcacggcaga   480

caaacaaaag aatggaatca aagctaactt caaaattcgc cacaacattg aagatggatc   540

cgttcaacta gcagaccatt atcaacaaaa tactccaatt ggcgatggcc ctgtcctttt   600

accagacaac cattacctgt cgacacaatc tgccctttcg aaagatccca acgaaaagcg   660

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gctctacaaa taa   733

<210> 2
<211> 6029
<212> DNA
<213> Artificial sequence

<220>
<223> Empty PETM-11 plasmid

<220>
<221> misc_feature
<222> (1)..(6029)
<223> Empty PETM-11 plasmid

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taaatctcgc gccgatcaac tgggtgccag cgtggtggtg tcgatggtag aacgaagcgg      1740

cgtcgaagcc tgtaaagcgg cggtgcacaa tcttctcgcg caacgcgtca gtgggctgat      1800

cattaactat ccgctggatg accaggatgc cattgctgtg gaagctgcct gcactaatgt      1860

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gctgttagcg ggcccattaa gttctgtctc ggcgcgtctg cgtctggctg gctggcataa      2040

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gtccggtttt caacaaacca tgcaaatgct gaatgagggc atcgttccca ctgcgatgct      2160

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gcaccattat gttccggatc tgcatcgcag gatgctgctg gctaccctgt ggaacaccta      3120

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ggtagctctt gatccggcaa acaaaccacc gctggtagcg gtggtttttt tgtttgcaag      4500

cagcagatta cgcgcagaaa aaaaggatct caagaagatc ctttgatctt ttctacgggg      4560
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<210> 3
<211> 5369
<212> DNA
<213> Artificial sequence

<220>
<223> Intermediate pETM-adp plasmid, on way to pETM-GFP-Imm construct

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gtggactctt gttccaaact ggaacaacac tcaaccctat ctcggtctat tcttttgatt   660

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attcgtgatt gcgcctgagc gagacgaaat acgcgatcgc tgttaaaagg acaattacaa   1260

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gaatcaggat attcttctaa tacctggaat gctgttttcc cggggatcgc agtggtgagt   1380

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agcgcttcgt taatacagat gtaggtgttc cacagggtag ccagcagcat cctgcgatgc   3240

agatccggaa cataatggtg cagggcgctg acttccgcgt ttccagactt tacgaaacac   3300
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